## DESCRIPTION OF NATIONAL GENETIC EVALUATION SYSTEMS

Country (or countries)	BELGIUM (Walloon Region)				
Country (or countries)					
Main trait group <sup>1</sup> NOTE! Only one trait group per form!	Birth Traits				
Breed(s)	Belgian Blue				
Trait definition(s) and unit(s) of	- Gestation length (days)				
measurement <sup>2</sup>	- Birth weight (kg)				
Attach an appendix if needed	- Body conformation score (1 - 9)				
Method of measuring and collecting data	By technicians or breeders				
Time period for data inclusion	All available data since 1993				
Age groups (e.g. parities) included	All				
Other criteria (data edits) for inclusion of	25 < birth weight < 70				
records	264 < gestation length < 301				
	0 < age at recording < 120 N/A				
<b>Criteria for extension of records</b> (if applicable)	NA				
Sire categories	All				
Environmental effects <sup>3</sup> , pre-adjustments	No pre-adjustments				
Method (model) of genetic evaluation <sup>3</sup>	Multiple trait animal model				
Environmental effects <sup>3</sup> in the genetic	Herd (F)				
evaluation model	Parity (F)				
	Year and Month of birth (F)				
	Sex (F) Suckling or not				
	Suchang of not				
Adjustment for heterogeneous variance in	No Adjustment				
evaluation model					
Use of genetic groups and relationships	No				
Blending of foreign/Interbull information	No blending				
in evaluation					
Genetic parameters in the evaluation	See Appendix GE				
System validation	Genetic trends, correlations between consecutive evaluations				
Expression of genetic evaluations	standardized breeding values, which are multiplied by a				
If standardised (e.g. RBV), give	standard error of 10 and added to a mean value of 100				
standardisation formula in the appendix					
Definition of genetic reference base	All cows born 7 years before the current year				
Next base change	Base changes every year				
Calculation of reliability	Reliabilities are calculated from PEV				
Criteria for official publication of	Sires : REL $\geq$ 50 % ; At least 10 calves in minimum 5 herds				
evaluations	Females : $REL \ge 15 \%$				
Number of evaluations / publications per	3				
year Use in total manifindar <sup>4</sup>	Ne				
Use in total merit index <sup>4</sup>	No				
Anticipated changes in the near future	/				
Key reference on methodology applied					

Key reference on methodology applied

Key organisation: name, address, phone, fax, e-mail, web site	Organisation responsible for genetic evaluations and computing centre:		
	Elevéo asbl		
	R&D Department - Genetic Evaluation Unit		
	Rue des Champs Elysées 4		
	B-5590 Ciney		
	0032/83/23.06.32.		
	eval_gen@awenet.be		
	WEB site for publication of sire breeding values: http://www.awenet.be		

1) Either: Production (e.g. milk, fat, protein), Conformation, Health (e.g. mastitis resistance, milk somatic cell, resistance to diseases other than mastitis), Longevity, Calving (e.g. stillbirth, calving ease), Female fertility (e.g. non-return rate, interval between reproductive events, number of AI's, heat strength), Workability (e.g. milking speed, temperament), Beef production, Efficiency (e.g. body weight, energy balance, body conditioning score), or Other traits.

2) Indicate frequencies per category if the trait is categorical and specify transformation of data if practiced.

3) Use abbreviations for most common effects (see document with list of abbreviations at http://www-

interbull.slu.se/service\_documentation/General/list\_of\_abbreviations.rtf) and indicate random (R) or fixed (F).

4) Please give economic weights and indicate how they are expressed (preferably in genetic standard deviation units).

## Parameters used in genetic evaluation

Country (or countries):	BELGIUM (Walloon Region)		
Main trait group: Breed (repeat as necessary):	Birth Traits Belgian-Blue		
		constis	official proc

Trait	Definition	ITB <sup>a</sup>	h <sup>2b</sup>	genetic variance <sup>b</sup>	official proof standardisation formula <sup>c</sup>
Gestation length			0.16	2,1	
Birth weight			0.09	5,3	
Conformation score			0.03	0,06	

<sup>a</sup> Indicate, with X, traits that are submitted to Interbull for international genetic evaluations.

<sup>b</sup> If repeated records are treated as separate traits, provide heritability estimates and genetic variances separately for each trait, as well as for all traits pooled, i.e. for the trait submitted to Interbull.

<sup>c</sup> Expressed as follows:

StandEval=((eval-a)/b)\*c+d where a=mean of the base adjustment, b=standard deviation of the base, c=standard deviation of expression (include sign if scale is reversed), and d=base of expression.